

SEQUENCE LISTING

<110> Ruvkun, Gary  
Ashrafi, Kaveh

<120> Polynucleotide and Polypeptide Fat  
Metabolism REGULATORS and Uses Thereof

<130> 00786/423002

<150> 60/395,159

<151> 2002-07-11

<160> 12

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 5570

<212> DNA

<213> *Caenorhabditis elegans*

<400> 1

agccctgttgc ctctgtcgaa gccgtttgt ctctctctcg tcgtcgctcg cccaccggc 60  
ctcccatgtt gtgtcgcttt tgcgtcttat cgcacacaca cactctcggt aaacaacgac 120  
cacctcactc catctctatc cattctatcc tttccatctt ctacaggcca acggagacag 180  
ttcggttggac ccccgccac tctacctatc tgttcaacgc tccatatgtg caaagtctca 240  
gtcatttttc ctccctgttc ttcttctttt tattataac ttttctcat cttatttctt 300  
ccactcttc gagagaccac tccgcccact ttgcgttact gctttatcaa acaaactgtg 360  
cgtccacacc aattctctat ttcccttcgt ctgctccgtt ttgcgttact ttttgttaa 420  
ctatctttc tttttcttt taacatgtt attgatcctc accactgtc aattaatatt 480  
tattttat ttctgttcc agatcaccta cgaaaaatata taattgataa tcagaggagt 540  
aaaaatgagg acatgtctca ccctcacggg tttcctctg atttcaatgg ccaccatttc 600  
ggggggcctc caaccgatgg ggcacactac aagaagtatg ttttctttt catttaataa 660  
tcacatagta attcaattga aaatcaatgc atttcgaat ttgcacat tactttgtt 720  
tgaccgttg agtcaccaat agatcaccta cgaaaaatata taattgataa tcagaggagt 780  
aaaaatgagg acatgtctca ccctcacggg tttcctctg atttcaatgg ccaccatttc 840  
ggggggcctc caaccgatgg ggcacactac aagaagtatg ttttctttt catttaataa 900  
tcacatagta attcaattga aaatcaatgc atttcgaat ttgcacat tactttgtt 960  
tgaccgttg agtcaccaat atcacgagat aataatttt tgaacaatgc caattttgtt 1020  
tcagaatgcg acgcaacaaa ttgcgttccag tgtcaagatg gccgatgc accgatgtcg 1080  
tggcgttgc atggagatat cgactgccag aatgaagaag atgaaaaaaa ttgtccaagt 1140  
aaaaactcttt ttctaaaaaa aacaatgata atttcaatga tagttcaga agtttgcgc 1200  
gccaagaac acaaatgcgg agaagtcaaa ttcgtctcgat catcggttga aagattcaag 1260  
tgcattccga acaaatgggt gtgtgatgg gaattcgact gtgaagataa atcggacgaa 1320  
ttccaatgca aaagtatgtt caattgaatt caagacagtt ttccgtcaat ttttcaatct 1380  
tttcagacgt atcatgccaa gaaaaacagt ttcaatgtga agaactctct ggtgattata 1440  
gtttgtgcat ttctgaaaca tgggttgcg atggtaaag agactgtacg aatggcaaag 1500  
acgaacagaa ttgcacgtca aaaacgtcta aatgtccgga taacaacttc cagtaagtct 1560  
ccatacaaat ttatcgcaa ttaaatctat cccgttgcg taattgtacc actctgacaa 1620  
aatttggaaaa aaagtgttcc atttttcagg ttttagcaatg gaaattgtat ttcaaaaaac 1680  
tgggttgcg atggggaaaga agattgtca gatggctcg atgaattgtct cactgctcca 1740  
tccaattgtt accgaacagt taatcaatgc cttccggag aaatgtggaa ggtatgatgt 1800  
tttacggatc ttgagaaaaat gtttatgcgc agcatgttga aatattttt tttttctcga 1860  
ttgttttagag ggtcaaagta tcgagtgtt gctatgttag tttcaaaacca actgataata 1920  
agaaaaatattt attatattga taattcattt aatatctatt tttctcataa tataagacttt 1980  
attaaataca taatacatac tagaaaaagaa aaataattca ataagctgga gtaagaaatt 2040

atcattttca gtcttttctt tgtcaattga ttgtgtattg cagtcgattg ttatctgtta 2100  
gtgaccattt ctcccccgca atcacgtctg ggaattatgtat atgtatttct gaattaaaca 2160  
attaaatttt cagtgtggat ccggtaatgc cattccatca agatggcgat gcgacgcaga 2220  
agtcgattgc aaagatcatt ccgatgagaa gaattgtact gctattcaac acacgtcaa 2280  
gttagcagag gtaggatggg ggcaatatgc acctgttgc tgctaatgca acatggctat 2340  
taaaatatta gatgtcaccc atattagata ggctaaattgg gttttaatc attaaatgtat 2400  
gagaattta acaacccat tatagttcca aaatgcaata acttcaggaa ttgcgttgc 2460  
aagcttcaca caactgcata aacaaggctt tcgtatgtga tggagaactt gattgttccg 2520  
atggatccga cgaagatgat tgcgtgcacg ttccgaccga gtgtaaatcc ggagagcgt 2580  
cctgcccagc ttcatacggc gcatatggcg ccgagtcagg tcacgttgc tgcgttgc 2640  
catcgatcg gtcaatggg gaagaggatt gtccagatgg tggtgatgag aaagaatgt 2700  
atatgactgc tcctggtaag taatattac aaaagtgcct gaaatataatt cgttatattt 2760  
tcagtcacat gccagaaaagg aaccgaatat gaatgtccat ctactccatt gcaatgtatt 2820  
gaaatgtcaa aatttgtgc tagtgcctaa tttgattgtg gggatggaaa tatgtctgtt 2880  
tgcgtccaga aaaagatcat tggtaatata aatattttat tatggtctgg caaaatgctc 2940  
ttcaattaca gaaatgtca aaccaagttc agaaggatgt gtctgcgtc catcgtttgt 3000  
ccgaggaat aatgttgc tttccatggc tttttttttt tttttttttt tttttttttt 3060  
tggtgagta ttgttagcaa gagagatggt ttagcgagaa aacaaatagt gaaaaagaca 3120  
aagagatcct catcaaaaatg tagaaaaata gttgagatgc gaagcggaca gctgaacaat 3180  
cagcaatatg aaaacacagg aagtatttc taataacgaa atgtttattt ttcagatatt 3240  
aacgaatgtg aaattgtcgg cggtgtgtt gtcatgtcg atatcagatt agttcggag atactaaaat tggatcagga 3360  
agaattgcta acaaatgtcg tgcgtggg ggtgatccat tggttcttct taccacacaga 3420  
catacgatca gacaatttgc tcttgcataat aaaatgcact tccctgtttc cagtagtcct 3480  
ggttctgcgg ttgcctatggg tttccacatc ttgaacgggg tgagttgaat tttttttttt 3540  
ggattgttat tatttacttg agaagaatac caaaaaatct gattttataa taattttttt 3600  
agacactgtat ttggctgtac gtgtgtcata agcaatttct gaaatgttcg attggaaacg 3660  
tgtcaaacgc atttttggg accgatatgt gcgataagaa gcatgaaatc gttctcacag 3720  
gagacaagat tcatactcca gatggacttgc cagttgatttgc gtttcatgat ctccctttct 3780  
ggacagatgg aggcttgc tttttttttt tttttttttt tttttttttt tttttttttt 3840  
tcctttattt ttccgacttgc caaattaatg ttctcgatat gaaaaatggaa aagcagcgt 3900  
tcattttctg gactgattgg gaagaaccga gagtatttgc agttgatcct gaagttggac 3960  
aacatcgatc tggttattgtt tttttttttt tttttttttt tttttttttt tttttttttt 4020  
attatgttgc taagcggtc tattggctga tgccaaatgc aagtcaatct tcagttgtga 4080  
ttattttttttt aaaaatatca agaccgtatttgc gcatctcat caatatctaa ggcattccatt 4140  
ctcaatggct gtttgcgaag accgacttctt ctacacagat tgggagcatg atggtgttat 4200  
cactgttaac aaggatgttt tttttttttt tttttttttt tttttttttt tttttttttt 4260  
aacgaaattt gctgaaaattt tttttttttt tttttttttt tttttttttt tttttttttt 4320  
atgaaaattt tctacacggaa atgaattttat tttttttttt tttttttttt tttttttttt 4380  
taaatcaatc ggcgtttcaa ctacgttgc tttttttttt tttttttttt tttttttttt 4440  
actgttatgg atcaagtggaa gtctccaatg actgtacgca tctaccacaa acaagcacaa 4500  
ccacttatgc agaacaagtg tggaaactcg gaatgtgatc atctctgcct gccgagagcc 4560  
gtttaccgtg agaaagaacg tggtaatgc tttttttttt tttttttttt tttttttttt 4620  
gcatgcgaag gaacgactgc tttttttttt tttttttttt tttttttttt tttttttttt 4680  
gtattttagat acttaacgtt tttttttttt tttttttttt tttttttttt tttttttttt 4740  
gatgttgcacg attttccttc tttttttttt tttttttttt tttttttttt tttttttttt 4800  
tgttcgctgg aagatgggac ctctgttgc tttttttttt tttttttttt tttttttttt 4860  
tcgtcgaacc accgaagaag ctgtatcatca gatggaagat ccattccgtg atccttttgc 4920  
tgaaccacgg aatggaagag ggcgttaacga tggattacca actcttgcat ctgctgacaa 4980  
tgaacacacgg gctgacgcatt tgagcttgc agccaattcg tattgtaaag tttttttttt 5040  
cctataaatt tatttgcacc tttttttttt tttttttttt tttttttttt tttttttttt 5100  
tcatgtcact tttttttttt gttttttttt ttctgttgc tttttttttt tttttttttt 5160  
ttcatcaatt tccattttttt gttttttttt gttttttttt tttttttttt tttttttttt 5220  
gtgaagacgt gtagatccaa actgtgaaaa tttttttttt tttttttttt tttttttttt 5280  
tatctatcc gaatggtttgc gatagtattt tgcacgttgc tttttttttt tttttttttt 5340  
gctttcatta ttctcattgc tatatcatta tttttttttt gttttttttt tttttttttt 5400  
ttatcgagct tttttttttt gttttttttt tttttttttt tttttttttt tttttttttt 5460  
tggtaattttt gttttttttt gttttttttt tttttttttt tttttttttt tttttttttt 5520  
atctgccccct tccctctcat caacggattt attcaataaa tttttttttt tttttttttt 5570

<210> 2  
<211> 2592  
<212> DNA  
<213> *Caenorhabditis elegans*

<400> 2  
atgaggacat gtctcaccct cacgggtttc cttctgattt caatggccac catttcggtg 60  
ggcctccaaac cgatgggagc acctacaaga aaatgcgacg caacaaaattc gttccagtgt 120  
caagatggcc gatgcatacc gatgtcgtgg cgttgtgatg gagatatcga ctgccagaat 180  
gaagaagatg agaaaaattt tccaaaagtt tttggcccg aagaacacaa atgcggagaa 240  
gtcaaatctg ctcgatcatc gttggaaaga ttcaagtgc ttccgaacaa atgggtgtgt 300  
gtatggagaat tcgactgtga agataaaatcg gacgaattcc aatgcacaaa cgtatcatgc 360  
caagaaaaac agtttcaatg tgaagaactc tctgggtatt atagtttgc cattccctgaa 420  
acatgggttt gcgatggtca aagagactgt acgaatggca aagacgaaca gaattgcacg 480  
tcaaaaacgt ctaaatgtcc ggataacaac ttccagtgta gcaatggaaa ttgtattttc 540  
aaaaactggg tttgtatgg ggaagaagat tgctcagatg gtcagatga attgctact 600  
gctccatcca attgtaaaccg aacagttaat caatggccctc ccggagaaaat gtggaagtgt 660  
ggatccgggtg aatgcattcc atcaagatgg cgatgcgacg cagaagtgcg ttgcaaaagat 720  
cattccgatg agaagaattt tactgttatt caacacacgt gcaagttgc agaggaattc 780  
gtttgttaaaag cttcacacaa ctgcacatcaac aaggcttcg tatgtatgg agaacttgat 840  
tggtccgatg gatccgacga agatgattgt gctgacgttc ggaccgagtg taaatccgga 900  
gagcgtaccc gcccagcttc atacgggtca tatggcccg agtcagggtca cgttgtgtgt 960  
attcctgcat cgtcatggtg caatggagaa gaggattgtc cagatggtg tgatgagaaa 1020  
gaatgtataa tgactgctcc tgcacatgc cagaaggaa cggaaatatga atgtccatct 1080  
actccattgc aatgtattga aatgtaaaaa ttgtgtgcta gtgctcaatt tgattgtggg 1140  
gatggaaataa tgcgttttgc tagccagaaa aagatcattt aatgtgcgca accaagttca 1200  
gaaggatgtg tctgcgtcc atcgtttgtc cgagggaaat atgttgtc ctgtaaagat 1260  
gttacaaac tcgaaaacgg acagtgcattt gatataacg aatgtgaaat tgctggcggtt 1320  
tgtatcaaa ttgtgtctaa tattcccggt tcctatcggtt gtgcttgcgatgat 1380  
cagattaggatc tcggagatataaatttggaa tcgagaaat ttgctaacaat atgtcggtct 1440  
atggggaggtt atccattgtt tcttcttacc aacagacata cgatcagacat atttgcattt 1500  
gtcaataaaa tgcacatccc ttgttcccgat agtccatgggtt ctgcgggttgc catggatttc 1560  
cacatcttgc acgggacact gatgggttgc gacgtgttgc caaagcaaat tctgaaatgt 1620  
tcgatggaa acgtgtcaaa cgcatttttggaaaccgata tgcgtatgcgatgaa 1680  
atcggtctca caggagacaa gattcataact ccagatggac ttgcgttgc ttgggttcat 1740  
gatctcattt tctggacaga tggaggcctt gatcaattt atgttctcgatgatgaaat 1800  
ggaaaaggcgc gtgtccttta ttcttccgac ttggaaagaaac cgagagctat tgcaatgtat 1860  
cctgaagttt gactcattttt ctggactgtat tggggaaagaaaggcgcgatgaaat cgaaagatct 1920  
ggaatggatg gacaacatcg tactgttatt ttggggagatcgatgtgtt atggccgaat 1980  
ggatggcgtt tggattatgt tgataaggcgat gtctattggc tgatgcacaa atcaagtca 2040  
tcttcgttt tcactggagc tgatattcgat actgttatgg atcaagtgcgatgaaatg 2100  
actgtacgca tctaccacaa acaagcacaa ccacttatgc agaacaatgt tgaaaactcg 2160  
gaatgtgatc atctctgcgtt gcccggagcc gtttaccgttgc agaaaaggaaat tgatgcatt 2220  
aaaaacttgc acgacagacc gttctcgat gcatgcgaaat gaaacgactgc ttctgtatgtt 2280  
ctggaaatgtt tcgctgactt gaaacaaaaa tccggaaatctt cgatgttgc gatttccctt 2340  
cttttatgtt ttgggtggat tgcgttgcgtt ggattttgttgc ttgttgcgttgc gaagatggaa 2400  
cctcgatcat ttaccgcctt caattttgcac aatccaaat tgcgtcgaac caccgaagaa 2460  
gtgtatcatc agatggaaaga tccattccgtt gatcccttttgcgttgc gaaaccacg gaatggaaaga 2520  
gggcgtaaacg atggattacc aactcttgcac tctgcgtaca atgaaacacg ggctgacgca 2580  
ttgagttctt ga 2592

<210> 3  
<211> 863  
<212> PRT  
<213> *Caenorhabditis elegans*

<400> 3

Met Arg Thr Cys Leu Thr Leu Thr Gly Phe Leu Leu Ile Ser Met Ala  
 1 5 10 15  
 Thr Ile Ser Val Gly Leu Gln Pro Met Gly Ala Pro Thr Arg Lys Cys  
 20 25 30  
 Asp Ala Thr Asn Ser Phe Gln Cys Gln Asp Gly Arg Cys Ile Pro Met  
 35 40 45  
 Ser Trp Arg Cys Asp Gly Asp Ile Asp Cys Gln Asn Glu Glu Asp Glu  
 50 55 60  
 Lys Asn Cys Pro Lys Val Cys Gly Ala Glu Glu His Lys Cys Gly Glu  
 65 70 75 80  
 Val Lys Ser Ala Arg Ser Ser Leu Glu Arg Phe Lys Cys Ile Pro Asn  
 85 90 95  
 Lys Trp Val Cys Asp Gly Glu Phe Asp Cys Glu Asp Lys Ser Asp Glu  
 100 105 110  
 Phe Gln Cys Lys Asn Val Ser Cys Gln Glu Lys Gln Phe Gln Cys Glu  
 115 120 125  
 Glu Leu Ser Gly Asp Tyr Ser Leu Cys Ile Pro Glu Thr Trp Val Cys  
 130 135 140  
 Asp Gly Gln Arg Asp Cys Thr Asn Gly Lys Asp Glu Gln Asn Cys Thr  
 145 150 155 160  
 Ser Lys Thr Ser Lys Cys Pro Asp Asn Asn Phe Gln Cys Ser Asn Gly  
 165 170 175  
 Asn Cys Ile Phe Lys Asn Trp Val Cys Asp Gly Glu Glu Asp Cys Ser  
 180 185 190  
 Asp Gly Ser Asp Glu Leu Leu Thr Ala Pro Ser Asn Cys Asn Arg Thr  
 195 200 205  
 Val Asn Gln Cys Pro Pro Gly Glu Met Trp Lys Cys Gly Ser Gly Glu  
 210 215 220  
 Cys Ile Pro Ser Arg Trp Arg Cys Asp Ala Glu Val Asp Cys Lys Asp  
 225 230 235 240  
 His Ser Asp Glu Lys Asn Cys Thr Ala Ile Gln His Thr Cys Lys Leu  
 245 250 255  
 Ala Glu Glu Phe Ala Cys Lys Ala Ser His Asn Cys Ile Asn Lys Ala  
 260 265 270  
 Phe Val Cys Asp Gly Glu Leu Asp Cys Ser Asp Gly Ser Asp Glu Asp  
 275 280 285  
 Asp Cys Ala Asp Val Arg Thr Glu Cys Lys Ser Gly Glu Arg Thr Cys  
 290 295 300  
 Pro Ala Ser Tyr Gly Ala Tyr Gly Ala Glu Ser Gly His Val Val Cys  
 305 310 315 320  
 Ile Pro Ala Ser Ser Trp Cys Asn Gly Glu Asp Cys Pro Asp Gly  
 325 330 335  
 Gly Asp Glu Lys Glu Cys Asn Met Thr Ala Pro Val Thr Cys Gln Lys  
 340 345 350  
 Gly Thr Glu Tyr Glu Cys Pro Ser Thr Pro Leu Gln Cys Ile Glu Met  
 355 360 365  
 Ser Lys Leu Cys Ala Ser Ala Gln Phe Asp Cys Gly Asp Gly Asn Met  
 370 375 380  
 Ser Val Cys Ser Gln Lys Ile Ile Glu Met Cys Lys Pro Ser Ser  
 385 390 395 400  
 Glu Gly Cys Val Cys Arg Pro Ser Phe Val Arg Gly Asn Asn Val Cys  
 405 410 415  
 His Cys Lys Asp Gly Tyr Lys Leu Glu Asn Gly Gln Cys Ile Asp Ile  
 420 425 430  
 Asn Glu Cys Glu Ile Ala Gly Val Cys Asp Gln Ile Cys Leu Asn Ile  
 435 440 445  
 Pro Gly Ser Tyr Arg Cys Ala Cys His Ala Gly Tyr Gln Ile Ser Phe  
 450 455 460  
 Gly Asp Thr Lys Ile Gly Ser Gly Arg Ile Ala Asn Lys Cys Arg Ala

465	470	475	480												
Met	Gly	Gly	Asp												
485	490	495													
Gln	Phe	Asp	Leu	Val	Asn	Lys	Met	His	Phe	Pro	Val	Ser	Ser	Ser	Pro
500	505	510													
Gly	Ser	Ala	Val	Ala	Met	Asp	Phe	His	Ile	Leu	Asn	Gly	Thr	Leu	Ile
515	520	525													
Trp	Ser	Asp	Val	Leu	Ser	Lys	Gln	Ile	Leu	Lys	Cys	Ser	Ile	Gly	Asn
530	535	540													
Val	Ser	Asn	Ala	Phe	Leu	Gly	Thr	Asp	Met	Cys	Asp	Lys	Lys	His	Glu
545	550	555	560												
Ile	Val	Leu	Thr	Gly	Asp	Lys	Ile	His	Thr	Pro	Asp	Gly	Leu	Ala	Val
565	570	575													
Asp	Trp	Val	His	Asp	Leu	Leu	Phe	Trp	Thr	Asp	Gly	Gly	Leu	Asp	Gln
580	585	590													
Ile	Asn	Val	Leu	Asp	Met	Lys	Asn	Gly	Lys	Gln	Arg	Val	Leu	Tyr	Ser
595	600	605													
Ser	Asp	Leu	Glu	Glu	Pro	Arg	Ala	Ile	Ala	Val	Asp	Pro	Glu	Val	Gly
610	615	620													
Leu	Ile	Phe	Trp	Thr	Asp	Trp	Gly	Lys	Lys	Ala	Arg	Ile	Glu	Arg	Ser
625	630	635	640												
Gly	Met	Asp	Gly	Gln	His	Arg	Thr	Val	Ile	Val	Glu	Gly	Asp	Arg	Val
645	650	655													
Val	Trp	Pro	Asn	Gly	Leu	Ala	Leu	Asp	Tyr	Val	Asp	Lys	Arg	Val	Tyr
660	665	670													
Trp	Leu	Met	Pro	Arg	Ser	Ser	Gln	Ser	Ser	Val	Phe	Thr	Gly	Ala	Asp
675	680	685													
Ile	Arg	Thr	Val	Met	Asp	Gln	Val	Lys	Ser	Pro	Met	Thr	Val	Arg	Ile
690	695	700													
Tyr	His	Lys	Gln	Ala	Gln	Pro	Leu	Met	Gln	Asn	Lys	Cys	Glu	Asn	Ser
705	710	715	720												
Glu	Cys	Asp	His	Leu	Cys	Leu	Pro	Arg	Ala	Val	Tyr	Arg	Glu	Lys	Glu
725	730	735													
Arg	Val	His	Glu	Lys	Thr	Trp	His	Asp	Arg	Pro	Phe	Ser	Cys	Ala	Cys
740	745	750													
Glu	Gly	Thr	Thr	Ala	Ser	Asp	Val	Leu	Glu	Cys	Phe	Ala	Asp	Leu	Glu
755	760	765													
Thr	Lys	Ser	Gly	Ile	Ser	Met	Phe	Thr	Ile	Phe	Leu	Leu	Cys	Val	
770	775	780													
Gly	Gly	Val	Val	Ala	Ala	Gly	Phe	Val	Ile	Val	Arg	Arg	Lys	Met	Gly
785	790	795	800												
Pro	Arg	Thr	Phe	Thr	Ala	Leu	Asn	Phe	Asp	Asn	Pro	Ile	Tyr	Arg	Arg
805	810	815													
Thr	Thr	Glu	Glu	Ala	Asp	His	Gln	Met	Glu	Asp	Pro	Phe	Arg	Asp	Pro
820	825	830													
Phe	Ala	Glu	Pro	Arg	Asn	Gly	Arg	Gly	Arg	Asn	Asp	Gly	Leu	Pro	Thr
835	840	845													
Leu	Ala	Ser	Ala	Asp	Asn	Glu	Thr	Arg	Ala	Asp	Ala	Leu	Ser	Phe	
850	855	860													

<210> 4

<211> 7495

<212> DNA

<213> *Caenorhabditis elegans*

<400> 4

tctccacttt caactggtca gagacgtcgt cttaacatc ttccccgtcg tcttccgcct 60  
aaaaaaagtgc gaaaagaaaac atcaacagaa aacaatgaat tgatcactac aattatataa 120

atttgccttt cttcctatca catatcaactt cgtctgtctg cgtctctatc actttattat 180  
 cttcaatatac ccacattatc tcgggttggcc tggaaacctt tcagtcgtt tttcttaaaa 240  
 ctattcatcg tcagcaacct cgtcatctt aaaaattaga aaaattggaa gaaaaaagag 300  
 aaataaaaaaa ggggtggagc ctagacacct tcaacacata ttttaatta aagacgccct 360  
 ttttcggaa gaccccttc tccgcttcc ccccattatt ttcttattttt atctaactga 420  
 tggaaaagccg aaaaacgag cccacttggg tgactaagcc tctgcttaaa cggatgagt 480  
 tgtcaagaga ttctctgaaa aaaacctaaa attttgaat attcaaaaaca gataatttca 540  
 gattcttagt atttgcata attccaaaat aaaaatata aacattttac agctctcatt 600  
 caagtgactc ttcaatcgat gaatcaactg ttaaactcac aaattatggg atattctatt 660  
 acactcaagg agttgatcta cttctttaa ttacttggaa acgttgcagca gttattcatg 720  
 gagctggttt tccgcttactt gctattgta ttttttaatt tgaatgataa 780  
 atcaaaaagct gaaattatca tttgaaacgt caactacata taatttata aaatgttatt 840  
 atgagaactc atagtcagaa ttaattttt ttttggaaa tttagttaaac tctaattctac 900  
 gttcaacatt cacaatataa cctccgatat cgtcatccaa tccaataaaac tactgcgact 960  
 acccattaaac ttaatttagat caaaaatgttcc atgacatcat ttgaactaga gaaaaaaagt 1020  
 gatttgcgtt gtggtttga actatggaaat tggaggctt ttatattctt caaaaaagga 1080  
 aatgtgtta agttgaaattt ttcagctcc ttaaaacaaa ttcaaaatatac atgagatatc 1140  
 ataggctgaa aattgtgata ttttataattt gcatatgtt gtatttaaa attttaaaaa 1200  
 tacactaact aagaagtcga acagatttca atatcataac taaacaatca aaaaatttct 1260  
 atagaaaaatg tggactttttt gagaattttt agaatttttgg cgggtttgaa gtcaagtttc 1320  
 cggaaaacaaa ataattaaaaa tataaaaactc gttagatatg ttttttagt tgacttccaa 1380  
 aattatgata aatcaaaaat aagggtatgg cacttttcg actgttgata agaaatttca 1440  
 aataatgtttt gaaaattttac attttggtca tttttttttt gatataatg ttatcataat ggttggttt 1500  
 aaataacttt ctcattaaac gaacatttctt gccccgcag taaaatcaat tagatagtt 1560  
 aaagcagaaaa cgtgatttcc aaagttctcg tatttgcagt ttcggagga atgacaacag 1620  
 tggttttacg agtcaaaaac tcggatttcg tcgttgggtt ggataatgtt aacccggaaag 1680  
 gattggtccc gatatctctg tacgtttttt ttagaactttt gacttttact tttttatgatc 1740  
 ctgcaattt ttgtattttt atcttttgcg ttagtcaatc tgcgcgaat gataaggcta 1800  
 tccaaaacag ctgggtgttc ttttgcacc ttttcttagt ctttgcgtt gaaacacataa 1860  
 atattatattt acattttttgcg attaaacaaa aaacatttcca atctgttttcc ttttttagag 1920  
 atgaatttca ctcggaaatgtt gtcaggatgtt gtagtacta cctgggttcc ggtgtactca 1980  
 tgttcttcac ttcatatgtt caaatcgatc gttttggatc gtacgcagag agattgggtgc 2040  
 ataaatatttac acaaaaacttca ctggaaatgtt tactcagaca acaatttcaat tgggtcgaca 2100  
 aacaacacagac cggaaaatgtt acggcttagac tcacggatgatgatgaaatgtt 2160  
 gaagaatgtt agagaatgtt gacatgtt atacatataa tgagtttttgcg cgggttgcgtca 2220  
 aatttttca gaaatttcatc taaattcccg gaagatcaat aagatatttgcg caaataataa 2280  
 atcatcgatc atcttttttgcg ttcggatgtt ccgttgcgtt ttaggtgaca 2340  
 aatttcggccct tcttgcgttca atgtttgcgtt ctttcttggc tggatgcgtt gttgggttct 2400  
 tttatagtttgcg ttcggatgttca atgtttgcgtt ctttcttggc tggatgcgtt gttgggttct 2460  
 gtgcggaaaat gggcaacgc gaaacatgtt gtcggatgtt ccgttgcgtt ttaggtgaca 2520  
 tcgctgtgtc aatttgcgttca gaaacatgtt ctttgcgtt aacatgttcat tcattttatg 2580  
 gacataaaatg agatgttgcgtt acggatgtt gtcggatgtt gttgggttct 2640  
 ttgtttaaata ttgttgcgtt ggttgcgtt gttggatgtt ttttgcgtt gttgggttct 2700  
 catatcgatc gggatgttgcgtt ttttgcgtt gttggatgtt ttttgcgtt gttgggttct 2760  
 gggatgttgcgtt ttttgcgtt gttggatgtt ttttgcgtt gttgggttct 2820  
 catatcgatc gggatgttgcgtt ttttgcgtt gttggatgtt ttttgcgtt gttgggttct 2880  
 ttttgcgtt gttggatgtt ttttgcgtt gttggatgtt ttttgcgtt gttgggttct 2940  
 ttttgcgtt gttggatgtt ttttgcgtt gttggatgtt ttttgcgtt gttgggttct 3000  
 gtttgcgtt gttggatgtt ttttgcgtt gttggatgtt ttttgcgtt gttgggttct 3060  
 gtttgcgtt gttggatgtt ttttgcgtt gttggatgtt ttttgcgtt gttgggttct 3120  
 gtttgcgtt gttggatgtt ttttgcgtt gttggatgtt ttttgcgtt gttgggttct 3180  
 gtttgcgtt gttggatgtt ttttgcgtt gttggatgtt ttttgcgtt gttgggttct 3240  
 gtttgcgtt gttggatgtt ttttgcgtt gttggatgtt ttttgcgtt gttgggttct 3300  
 gtttgcgtt gttggatgtt ttttgcgtt gttggatgtt ttttgcgtt gttgggttct 3360  
 gtttgcgtt gttggatgtt ttttgcgtt gttggatgtt ttttgcgtt gttgggttct 3420  
 gtttgcgtt gttggatgtt ttttgcgtt gttggatgtt ttttgcgtt gttgggttct 3480  
 gtttgcgtt gttggatgtt ttttgcgtt gttggatgtt ttttgcgtt gttgggttct 3540  
 gtttgcgtt gttggatgtt ttttgcgtt gttggatgtt ttttgcgtt gttgggttct 3600  
 gtttgcgtt gttggatgtt ttttgcgtt gttggatgtt ttttgcgtt gttgggttct 3660

aaaaataacg aaagtcatcg tttaaattta aaaaaaaatt atacatttac atatatttca 3720  
tattccagcc attgcacgtg ctctgtcaa aaatccaaaa atcctttgc tcgacgaagc 3780  
cacatccgct ctcgacacgg aagctgagag agaagttca ggtgcattgg atcagggcaca 3840  
agctggaga aacgacaatta tcgtagctca ccgattgagc acaattcgaa atgttgacag 3900  
aatattcgta ttcaaagctg gaaatattgt tgaatctgga agtcatgagg aattaatgag 3960  
caaacaagga atcttctacg atatgacaca ggctcaagtt gttcgacaac agcaacagg 4020  
agcaggaaaa ggttaattcta atgttaagg aaaactaata tagattaaat ttcatgatatt 4080  
gaagacacta ttcttgagtc agctcattcc catctcagca gaaagtcttc cacaagaagt 4140  
gccatttcaa ttgcaacatc tattcateag ctcgctgagg aggttgaggt acgaaaataa 4200  
ttacttattt ctttggttt taatagtac acttgaccaaa atataaaacc tctacaattt ttttagatatt ccattttgag 4260  
attaagagag tttgataaaa aagcaatgcc gcatgttgcg ttggcaaaatg tttgaaaaat tgggctttc aaagaaaattt 4320  
ttggcaaaatg tcttctacaa cggttatata caaattatca aaaaacacaa 4380  
tcttctacaa agaaaaattt ttagtcgac ttccaaaattt atgagttgcg 4440  
tattgactgt aaaaattaa tataattttt gaaaattttt 4500  
gaccataata tttgatgtta attcttctac tggcctact 4560  
ccatccttta aaaaataata aatccaaac aattatagtt aaacgaatca agatgtgata 4620  
gatgaatata ttccggtttc ttcaatgttca ttcaacttct ctccttctc tgattcttcc 4680  
ttacatttc ttcaaacacg gcttcttca agtacttac agcatgctt tatattgttt 4740  
ttttggttca atgatcaatt tttttaaat ttttcttaat ttaacaaaat aactttcagg 4800  
aatgcaggc tccacccacc tcaatgttca aaatattca attcaacgga gacaaagtgc 5040  
gatggtttat tggtggatt tttggagcat ttattttgg atcagttact ccagttttt 5100  
ctcttgata tgctgaaatt ctgtacattt ttcaatgttca ttcaatgttca ttcaacttct ctccttctc tgattcttcc 5160  
ttttttaaat ttttcttaat ttaacaaaat aactttcagg 5220  
tatgggaatc actttcttcg ttggatttctt cacttctgca 5280  
agagtcactg acaatgaagt tgagatttga agcattcaag 5340  
cgctttttat gatgatttga gacatggaa tggaaaattt 5400  
ctcttgcgca gctgatcaaa tgcaagcaaa tgtgtatttc 5460  
tatgggaatc actttcttcg ttggatttctt cacttctgca 5520  
agagtcactg acaatgaagt tgagatttga agcattcaag 5580  
cgctttttat gatgatttga gacatggaa tggaaaattt 5640  
ctcttgcgca gctgatcaaa tgcaagcaaa tgtgtatttc 5700  
tatgggaatc actttcttcg ttggatttctt cacttctgca 5760  
tgcatcttctt caatcttta ttcttcttcat gtatgtct 5820  
ttttgttaat caacaagcta tgcaaccaat tgatgtctat 5880  
attctgttgcg caaatgattt gaaataactac atctttttt 5940  
tcttgcgtct tctctttgt tctatctt tgaacatcca 6000  
tgatagttga attgtgaagc cgataactgg aatatttca 6060  
ttatccaaaca agaaaggata ccaagggtt acaaggattc 6120  
tttgataacctt gacttctata tgacagtagt gcaatcctag 6180  
gacttttaaa aactggatattt ggatttttt tgcgtttttt 6240  
gcccctgtgc tttttactt gaaattttaaa aataggaaaa 6300  
acaatttta acccacctgt aaaaacaaat attaatat 6360  
ttattttaga tcaaagccgg taaaactgtt gcacttgcg ggcactcagg atgtggaaaa 6420  
tctacaatta tggactgtc ggagagattt tataatcaag ataaaggaaat gattgtgagt 6480  
caattttctt tctgttgggt tttaactgtca aacaattttt gatgattgtt ggtgataaca 6540  
tccgttaacctt aaacatcagt tcacttcgcg aacaaggatgtt tattgttaatg caagagccaa 6600  
cggttttgcg ttgcacaattt ggagaaaaataa tttgctacgg aacaaatcga aatgttacat 6660  
atcaagaaaat tggactgtc gccaaaatgg caaatattca caatttcattt cttagattgc 6720  
cagatgttagg gtgatattttt cataaaatcga aactcattttt aaaaatttca gggtttatgtat 6780  
actcatgtcg gagagaaaagg aactcaactt tcgggtggtc aaaaacaaag aattgcccatt 6840  
gcacgggcac ttgttcgatc tccttctgtt ttacttttttgg atgaagcaac tagtgcatttt 6900  
gatacggaaa gtggaaaagggtt tggtatggaaa aatattgaaa tagcaaatttgcg atcttgcgaa 6960  
atatcggtt attcactgtt tacagattgtt acaagaagca ttggacgcgg caaaacaagg 7020  
tcgcacgtgtt cttgttcatgg gatcgtcagt gagggtaaaaa ttgtggaaaaa gggaaacacat gacgagttga taaggaagag 7140  
tgaaaatataat cagaaattctt gtgaaacgcg gaggattgtc gaaagtcaat aattttaaataa 7200

tgttagat	tctcaaacac	gagtttacaa	actaattgc	atggagttc	atttttttaa	7260
tgttcaattg	aaacagctt	atatttaaaa	tttaaatatg	ctcatcaagt	aaaatttttta	7320
aaaaattttt	taaacccgta	ataattttt	tgtcatctag	gtactttgct	ttttcccaa	7380
atagccttc	cctccatctt	gtgtattttt	tgtgaattt	tttgaattgt	gataattatc	7440
tttqaattgt	qataattgtc	tttttqttt	ctttttaaa	tatattattt	accat	7495

<210> 5  
<211> 3798  
<212> DNA  
<213> *Caenorhabditis elegans*

<400> 5  
atggaaaagcc gaaaaaaacga gcccaacttgg gtgactaagc ctctgcttaa acgctctcat 60  
tcaagtgaact cttcaatcga tgaatcaact gttaaactca caaattatgg gatattctat 120  
tacactcaag gagttgatct acttcttta attactggaa cagttgcagc agttattcat 180  
ggagctggtt ttccgttact tgctattgtt ctcggaggaa tgacaacagt gtttttacga 240  
gctcaaactaact cgatttcgt cggtgggtg gataatgtga acccggaaagg attggcccg 300  
atatctctag atgaattcaa ctcggaaagg gtcaagtatt gtatctacta cctggttctt 360  
ggtgtactca tggttcttcac ttcatatgtt caaatcgctt gttttgagtc gtacgcagag 420  
agattgggtc ataaattaag acaaaaactac ctgaaaagcca tactcagaca acaaattcaa 480  
tgggtcgaca aacaacagac cgaaaattta acggctagac tcacggacga tttggagcgt 540  
gtccgtgaag gattaggtga caaattcggcc cttctgttc aaatgtttgc tgctttctt 600  
gctggatacg gagttggctt ctttatagt tggtcaatga cactggttat gatgggattt 660  
gctccgttga ttgtgctctc tggtgccaaa atgagcaaaa gcatggcaac gcgaaacaaga 720  
gttgaacaag aaacgtatgc agtcgctggt gcaattgcag aagaaaacatt ctcttcgatt 780  
agaacagttc attcattaaa tggacataaa agagaatttg atagattttta taacgcattt 840  
gaagttggaa gacaaaactgg aattgttaaa tattgttata tgggtattgg agttgggtt 900  
agtaatttgt gtatgtactc ttcatatgca ttggcattttt ggatggaaag tactctgatt 960  
atcaatgatc ctactttga tcgccccgtt attttacgg ttttcttcgc agttctctcg 1020  
ggttctacat ctctcggtgg cgcccttcca catctgcaaa gttttggaaac agctcgccgaa 1080  
gcagcttcaa cagattacg tgaatcaac tcgcacccaa aaatcgatcc atattcaactt 1140  
gaaggaatttcc tcgtggacaa tatgaagggaa gatatttcat tcaaagatgt tcatttccga 1200  
tatccatctc gaaaagatata tcatgttataa aaaggaattt ctctggaaact gaaagctggt 1260  
gataaaaatttgc ctgggtcggtt tcaagtgggt tggggaaaat caacaattgtt taatttactt 1320  
caaagattct atgatccaac aaaaggaaga gttttaatttgc atggagttga ttacgagaa 1380  
gtaaatgttc atagtcttcg tgaacaaattt ggaattgttta gtcaagagcc agtactttc 1440  
gatggaaacaa ttatgaaaaa tattaaaatg ggaattgagc atgctactca tgatcaagtc 1500  
gttgaagcggtt gtaaaaatggc aaatgcaaaat gattttatca aaagattgccc tgatggat 1560  
ggaacaagag ttggagaaaaa aggagttcaa ttaagtggag gacagaaaaca aagaattgccc 1620  
attgcacgtt ctctgttcaa aaatccaaaaa atcctttgc tcgacgaaagc cacatccgct 1680  
ctcgacacccgaa aagctgagag agaagttcaa ggtgcattgg atcaggcaca agctggaaaga 1740  
acgacaatttca tcgtagctca ccgattgagc acaattcgaa atgttgcacag aatattcggt 1800  
ttcaaaagctt gaaatattgt tgaatctggaa agtcatgagg aattaatgag caaacaagga 1860  
atcttctacg atatgacaca ggctcaagggtt gttcgacaac agcaacagggaa agcagggaaaa 1920  
gatattgttgc acactatttc tgagtcaactt cattcccatc tcagcagaaaa gtcttcacaca 1980  
agaagtgccaa ttcaatttgc aacatctattt catcagctcg ctgaggaggt tgaggaatgc 2040  
aaggctccac ccacccat tttcaaaaatttca acggagacaa agtcggatgg 2100  
tttattgggtt gaaatttggt agcattttt gttggatcag ttactccagt ttttgcctt 2160  
gtatattgttgc aaattttcaaa ttttgcactt ttgcacgtt atcaaatgtca agcaaatgtg 2220  
tattttctgtt gttggatgtt ttttgcattt ggaatcactt ttttgcgttgg attcttcact 2280  
tctgcatttgc gtttttttttgcactt ttttgcattt ggaatcactt ttttgcgttgg attcttcact 2340  
ttcaagaatttataaagaca agatatcgct ttttgcattt gtttttttttgcactt ttttgcgttgg attcttcact 2400  
aaatttgcacaa caagatttgc aactgtatgtt ccgaatgttca gatatgttatttgcactt ttttgcgttgg attcttcact 2460  
ccagttgtttt tagcatcaat ttttgcattt gtttttttttgcactt ttttgcgttgg attcttcact 2520  
ggatggcaac ttgccttgcattt ttttgcattt ggaatcactt ttttgcgttgg attcttcact 2580  
ttcgaaatgc aaatgttgcattt ttttgcattt gtttttttttgcactt ttttgcgttgg attcttcact 2640  
ggaaaatgttgcattt ttttgcattt gtttttttttgcactt ttttgcgttgg attcttcact 2700

gaacaatttc attcacata ctgtaatat ctccggAAC cattcaatac taatctgaaa 2760  
 catgcacata catatggAGC tggatTTGCA ttctctcaat ctcttatttt ctcatgtat 2820  
 gctgctgcat tctatcttgg aagtattttt gtaaatcaac aagctatgca accaattgt 2880  
 gtcatacgag tattcttgc tatttcattc tggtggacaaa tgattggaaa tactacatct 2940  
 tttattcctg atgtcgtaaa agctcgctt gctgcttctc tttgttcta tcttattgaa 3000  
 catccaacac ctattgattc tctatctgat agtggaaattg tgaagccgat aactggaaat 3060  
 atttcaatca gaaatgtatt tttcaattat ccaacaagaa aggataccaa ggtttacaa 3120  
 ggattcactc ttgatataaa agccgtaaa actgttgcac ttgtcggc ctcaggatgt 3180  
 ggaaaatcta caattatggg actgctggag agattctata atcaagataa aggaatgatt 3240  
 atgattgatg gtgataacat ccgtaaccta aacatcagtt cacttcgcga acaagtatgt 3300  
 attgtaagtc aagagccaaac gttgttgc tgcacaattt gagaaaatat ttgtcgtcga 3360  
 acaaattcgaa atgttacata tcaagaaattt gtgaaatgtc ccaaaatggc aaatattcac 3420  
 aatttcatc taggattgcc agatgatattt gatactcatg tcggagagaa aggaactcaa 3480  
 cttcgggtg gtcaaaaaca aagaatttgc attgcacggg cacttgcgc atctccttct 3540  
 gtttacttt tggatgaagc aactagtgc ttgatatacgaa aagtggaaa gattgtacaa 3600  
 gaagcattgg acgcccacaa acaaggtcgc acgtgttgc tcattgtc tcgggttgc 3660  
 acaattcaaa atagtgacgt cattgcgatc gtcagtggg gtaaaattgt ggaaaaggga 3720  
 acacatgacg agttgataag gaagagtgaa atatatcaga aattctgtga aacgcagagg 3780  
 attgtcgaaa gtcaataaa 3798

<210> 6  
 <211> 1265  
 <212> PRT  
 <213> *Caenorhabditis elegans*

<400> 6  
 Met Lys Ser Arg Lys Asn Glu Pro Thr Trp Val Thr Lys Pro Leu Leu  
 1 5 10 15  
 Lys Arg Ser His Ser Ser Asp Ser Ser Ile Asp Glu Ser Thr Val Lys  
 20 25 30  
 Leu Thr Asn Tyr Gly Ile Phe Tyr Tyr Thr Gln Gly Val Asp Leu Leu  
 35 40 45  
 Leu Leu Ile Thr Gly Thr Val Ala Ala Val Ile His Gly Ala Gly Phe  
 50 55 60  
 Pro Leu Leu Ala Ile Val Leu Gly Gly Met Thr Thr Val Phe Leu Arg  
 65 70 75 80  
 Ala Gln Asn Ser Asp Phe Val Val Gly Val Asp Asn Val Asn Pro Glu  
 85 90 95  
 Gly Leu Val Pro Ile Ser Leu Asp Glu Phe Asn Ser Glu Val Val Lys  
 100 105 110  
 Tyr Cys Ile Tyr Tyr Leu Val Leu Gly Val Leu Met Phe Phe Thr Ser  
 115 120 125  
 Tyr Val Gln Ile Ala Cys Phe Glu Ser Tyr Ala Glu Arg Leu Val His  
 130 135 140  
 Lys Leu Arg Gln Asn Tyr Leu Lys Ala Ile Leu Arg Gln Gln Ile Gln  
 145 150 155 160  
 Trp Phe Asp Lys Gln Gln Thr Gly Asn Leu Thr Ala Arg Leu Thr Asp  
 165 170 175  
 Asp Leu Glu Arg Val Arg Glu Gly Leu Gly Asp Lys Phe Ala Leu Leu  
 180 185 190  
 Val Gln Met Phe Ala Ala Phe Leu Ala Gly Tyr Gly Val Gly Phe Phe  
 195 200 205  
 Tyr Ser Trp Ser Met Thr Leu Val Met Met Gly Phe Ala Pro Leu Ile  
 210 215 220  
 Val Leu Ser Gly Ala Lys Met Ser Lys Ser Met Ala Thr Arg Thr Arg  
 225 230 235 240  
 Val Glu Gln Glu Thr Tyr Ala Val Ala Gly Ala Ile Ala Glu Glu Thr  
 245 250 255  
 Phe Ser Ser Ile Arg Thr Val His Ser Leu Asn Gly His Lys Arg Glu

260	265	270
Leu	Asp	Arg
Phe	Tyr	Asn
275	280	285
Ala	Leu	Glu
Gly	Val	Gly
Arg	Arg	Gln
290	295	300
Tyr	Cys	Tyr
Met	Gly	Ile
305	310	315
Tyr	Ser	Ser
Tyr	Ala	Leu
320	325	330
Leu	Ala	Phe
335	340	345
Asn	Asp	Thr
345	350	355
Phe	Phe	Arg
355	360	365
365	370	375
380	385	390
395	400	405
410	415	420
425	430	435
440	445	450
455	460	465
470	475	480
485	490	495
500	505	510
515	520	525
530	535	540
545	550	555
560	565	570
575	580	585
590	595	600
605	610	615
620	625	630
640	645	650
655	660	665
670	675	680
685	690	695
700	705	710
715	720	725
730	735	

Gln Ala Asn Val Tyr Phe Trp Cys Gly Met Phe Val Leu Met Gly Ile  
 740 745 750  
 Thr Phe Phe Val Gly Phe Phe Thr Ser Ala Asn Cys Leu Gly Arg Cys  
 755 760 765  
 Gly Glu Ser Leu Thr Met Lys Leu Arg Phe Glu Ala Phe Lys Asn Leu  
 770 775 780  
 Leu Arg Gln Asp Ile Ala Phe Tyr Asp Asp Leu Arg His Gly Thr Gly  
 785 790 795 800  
 Lys Leu Cys Thr Arg Phe Ala Thr Asp Ala Pro Asn Val Arg Tyr Val  
 805 810 815  
 Phe Thr Arg Leu Pro Val Val Leu Ala Ser Ile Val Thr Ile Cys Gly  
 820 825 830  
 Ala Leu Gly Ile Gly Phe Tyr Tyr Gly Trp Gln Leu Ala Leu Ile Leu  
 835 840 845  
 Val Val Met Val Pro Leu Leu Val Met Gly Gly Tyr Phe Glu Met Gln  
 850 855 860  
 Met Arg Phe Gly Lys Gln Ile Arg Asp Thr Gln Leu Leu Glu Glu Ala  
 865 870 875 880  
 Gly Lys Val Ala Ser Gln Ala Val Glu His Ile Arg Thr Val His Ser  
 885 890 895  
 Leu Asn Arg Gln Glu Gln Phe His Phe Thr Tyr Cys Glu Tyr Leu Arg  
 900 905 910  
 Glu Pro Phe Asn Thr Asn Leu Lys His Ala His Thr Tyr Gly Ala Val  
 915 920 925  
 Phe Ala Phe Ser Gln Ser Leu Ile Phe Phe Met Tyr Ala Ala Ala Phe  
 930 935 940  
 Tyr Leu Gly Ser Ile Phe Val Asn Gln Gln Ala Met Gln Pro Ile Asp  
 945 950 955 960  
 Val Tyr Arg Val Phe Phe Ala Ile Ser Phe Cys Gly Gln Met Ile Gly  
 965 970 975  
 Asn Thr Thr Ser Phe Ile Pro Asp Val Val Lys Ala Arg Leu Ala Ala  
 980 985 990  
 Ser Leu Leu Phe Tyr Leu Ile Glu His Pro Thr Pro Ile Asp Ser Leu  
 995 1000 1005  
 Ser Asp Ser Gly Ile Val Lys Pro Ile Thr Gly Asn Ile Ser Ile Arg  
 1010 1015 1020  
 Asn Val Phe Phe Asn Tyr Pro Thr Arg Lys Asp Thr Lys Val Leu Gln  
 1025 1030 1035 1040  
 Gly Phe Thr Leu Asp Ile Lys Ala Gly Lys Thr Val Ala Leu Val Gly  
 1045 1050 1055  
 His Ser Gly Cys Gly Lys Ser Thr Ile Met Gly Leu Leu Glu Arg Phe  
 1060 1065 1070  
 Tyr Asn Gln Asp Lys Gly Met Ile Met Ile Asp Gly Asp Asn Ile Arg  
 1075 1080 1085  
 Asn Leu Asn Ile Ser Ser Leu Arg Glu Gln Val Cys Ile Val Ser Gln  
 1090 1095 1100  
 Glu Pro Thr Leu Phe Asp Cys Thr Ile Gly Glu Asn Ile Cys Tyr Gly  
 1105 1110 1115 1120  
 Thr Asn Arg Asn Val Thr Tyr Gln Glu Ile Val Glu Ala Ala Lys Met  
 1125 1130 1135  
 Ala Asn Ile His Asn Phe Ile Leu Gly Leu Pro Asp Gly Tyr Asp Thr  
 1140 1145 1150  
 His Val Gly Glu Lys Gly Thr Gln Leu Ser Gly Gly Gln Lys Gln Arg  
 1155 1160 1165  
 Ile Ala Ile Ala Arg Ala Leu Val Arg Ser Pro Ser Val Leu Leu Leu  
 1170 1175 1180  
 Asp Glu Ala Thr Ser Ala Leu Asp Thr Glu Ser Glu Lys Ile Val Gln  
 1185 1190 1195 1200  
 Glu Ala Leu Asp Ala Ala Lys Gln Gly Arg Thr Cys Leu Val Ile Ala

1205	1210	1215
His Arg Leu Ser Thr Ile Gln Asn Ser Asp Val Ile Ala Ile Val Ser		
1220	1225	1230
Glu Gly Lys Ile Val Glu Lys Gly Thr His Asp Glu Leu Ile Arg Lys		
1235	1240	1245
Ser Glu Ile Tyr Gln Lys Phe Cys Glu Thr Gln Arg Ile Val Glu Ser		
1250	1255	1260
Gln		
1265		

<210> 7  
 <211> 29  
 <212> DNA  
 <213> *Caenorhabditis elegans*

<400> 7		
cacaacaagt cagcaagcaa tacaagtgg		29

<210> 8  
 <211> 28  
 <212> DNA  
 <213> *Caenorhabditis elegans*

<400> 8		
gtaggagatg tgaccaatcg ttgaagtg		28

<210> 9  
 <211> 873  
 <212> PRT  
 <213> *Homo sapiens*

<400> 9			
Met Gly Thr Ser Ala Leu Trp Ala Leu Trp Leu Leu Leu Ala Leu Cys			
1	5	10	15
Trp Ala Pro Arg Glu Ser Gly Ala Thr Gly Thr Gly Arg Lys Ala Lys			
20	25	30	
Cys Glu Pro Ser Gln Phe Gln Cys Thr Asn Gly Arg Cys Ile Thr Leu			
35	40	45	
Leu Trp Lys Cys Asp Gly Asp Glu Asp Cys Val Asp Gly Ser Asp Glu			
50	55	60	
Lys Asn Cys Val Lys Lys Thr Cys Ala Glu Ser Asp Phe Val Cys Asn			
65	70	75	80
Asn Gly Gln Cys Val Pro Ser Arg Trp Lys Cys Asp Gly Asp Pro Asp			
65	90	95	
Cys Glu Asp Gly Ser Asp Glu Ser Pro Glu Gln Cys His Met Arg Thr			
100	105	110	
Cys Arg Ile His Glu Ile Ser Cys Gly Ala His Ser Thr Gln Cys Ile			
115	120	125	
Pro Val Ser Trp Arg Cys Asp Gly Glu Asn Asp Cys Asp Ser Gly Glu			
130	135	140	
Asp Glu Glu Asn Cys Gly Asn Ile Thr Cys Ser Pro Asp Glu Phe Thr			
145	150	155	160
Cys Ser Ser Gly Arg Cys Ile Ser Arg Asn Phe Val Cys Asn Gly Gln			
165	170	175	
Asp Asp Cys Ser Asp Gly Ser Asp Glu Leu Asp Cys Ala Pro Pro Thr			
180	185	190	
Cys Gly Ala His Glu Phe Gln Cys Ser Thr Ser Ser Cys Ile Pro Ile			
195	200	205	

Ser Trp Val Cys Asp Asp Asp Ala Asp Cys Ser Asp Gln Ser Asp Glu  
 210 215 220  
 Ser Leu Glu Gln Cys Gly Arg Gln Pro Val Ile His Thr Lys Cys Pro  
 225 230 235 240  
 Ala Ser Glu Ile Gln Cys Gly Ser Gly Glu Cys Ile His Lys Lys Trp  
 245 250 255  
 Arg Cys Asp Gly Asp Pro Asp Cys Lys Asp Gly Ser Asp Glu Val Asn  
 260 265 270  
 Cys Pro Ser Arg Thr Cys Arg Pro Asp Gln Phe Glu Cys Glu Asp Gly  
 275 280 285  
 Ser Cys Ile His Gly Ser Arg Gln Cys Asn Gly Ile Arg Asp Cys Val  
 290 295 300  
 Asp Gly Ser Asp Glu Val Asn Cys Lys Asn Val Asn Gln Cys Leu Gly  
 305 310 315 320  
 Pro Gly Lys Phe Lys Cys Arg Ser Gly Glu Cys Ile Asp Ile Ser Lys  
 325 330 335  
 Val Cys Asn Gln Glu Gln Asp Cys Arg Asp Trp Ser Asp Glu Pro Leu  
 340 345 350  
 Lys Glu Cys His Ile Asn Glu Cys Leu Val Asn Asn Gly Gly Cys Ser  
 355 360 365  
 His Ile Cys Lys Asp Leu Val Ile Gly Tyr Glu Cys Asp Cys Ala Ala  
 370 375 380  
 Gly Phe Glu Leu Ile Asp Arg Lys Thr Cys Gly Asp Ile Asp Glu Cys  
 385 390 395 400  
 Gln Asn Pro Gly Ile Cys Ser Gln Ile Cys Ile Asn Leu Lys Gly Gly  
 405 410 415  
 Tyr Lys Cys Glu Cys Ser Arg Gly Tyr Gln Met Asp Leu Ala Thr Gly  
 420 425 430  
 Val Cys Lys Ala Val Gly Lys Glu Pro Ser Leu Ile Phe Thr Asn Arg  
 435 440 445  
 Arg Asp Ile Arg Lys Ile Gly Leu Glu Arg Lys Glu Tyr Ile Gln Leu  
 450 455 460  
 Val Glu Gln Leu Arg Asn Thr Val Ala Leu Asp Ala Asp Ile Ala Ala  
 465 470 475 480  
 Gln Lys Leu Phe Trp Ala Asp Leu Ser Gln Lys Ala Ile Phe Ser Ala  
 485 490 495  
 Ser Ile Asp Asp Lys Val Gly Arg His Val Lys Met Ile Asp Asn Val  
 500 505 510  
 Tyr Asn Pro Ala Ala Ile Ala Val Asp Trp Val Tyr Lys Thr Ile Tyr  
 515 520 525  
 Trp Thr Asp Ala Ala Ser Lys Thr Ile Ser Val Ala Thr Leu Asp Gly  
 530 535 540  
 Thr Lys Arg Lys Phe Leu Phe Asn Ser Asp Leu Arg Glu Pro Ala Ser  
 545 550 555 560  
 Ile Ala Val Asp Pro Leu Ser Gly Phe Val Tyr Trp Ser Asp Trp Gly  
 565 570 575  
 Glu Pro Ala Lys Ile Glu Lys Ala Gly Met Asn Gly Phe Asp Arg Arg  
 580 585 590  
 Pro Leu Val Thr Ala Asp Ile Gln Trp Pro Asn Gly Ile Thr Leu Asp  
 595 600 605  
 Leu Ile Lys Ser Arg Leu Tyr Trp Leu Asp Ser Lys Leu His Met Leu  
 610 615 620  
 Ser Ser Val Asp Leu Asn Gly Gln Asp Arg Arg Ile Val Leu Lys Ser  
 625 630 635 640  
 Leu Glu Phe Leu Ala His Pro Leu Ala Leu Thr Ile Phe Glu Asp Arg  
 645 650 655  
 Val Tyr Trp Ile Asp Gly Glu Asn Glu Ala Val Tyr Gly Ala Asn Lys  
 660 665 670  
 Phe Thr Gly Ser Glu Leu Ala Thr Leu Val Asn Asn Leu Asn Asp Ala

675	680	685
Gln Asp Ile Ile Val Tyr His Glu Leu Val Gln Pro Ser Gly Lys Asn		
690	695	700
Trp Cys Glu Glu Asp Met Glu Asn Gly Gly Cys Glu Tyr Leu Cys Leu		
705	710	715
Pro Ala Pro Gln Ile Asn Asp His Ser Pro Lys Tyr Thr Cys Ser Cys		
725	730	735
Pro Ser Gly Tyr Asn Val Glu Glu Asn Gly Arg Asp Cys Gln Ser Thr		
740	745	750
Ala Thr Thr Val Thr Tyr Ser Glu Thr Lys Asp Thr Asn Thr Thr Glu		
755	760	765
Ile Ser Ala Thr Ser Gly Leu Val Pro Gly Gly Ile Asn Val Thr Thr		
770	775	780
Ala Val Ser Glu Val Ser Val Pro Pro Lys Gly Thr Ser Ala Ala Trp		
785	790	795
Ala Ile Leu Pro Leu Leu Leu Val Met Ala Ala Val Gly Gly Tyr		
805	810	815
Leu Met Trp Arg Asn Trp Gln His Lys Asn Met Lys Ser Met Asn Phe		
820	825	830
Asp Asn Pro Val Tyr Leu Lys Thr Thr Glu Glu Asp Leu Ser Ile Asp		
835	840	845
Ile Gly Arg His Ser Ala Ser Val Gly His Thr Tyr Pro Ala Ile Ser		
850	855	860
Val Val Ser Thr Asp Asp Asp Leu Ala		
865	870	

<210> 10  
 <211> 873  
 <212> PRT  
 <213> Rattus norvigicus

<400> 10		
Met Gly Thr Ser Ala Arg Trp Ala Leu Trp Leu Leu Leu Ala Leu Cys		
1	5	10
Trp Ala Pro Arg Asp Ser Gly Ala Thr Ala Ser Gly Lys Lys Ala Lys		
20	25	30
Cys Asp Ser Ser Gln Phe Gln Cys Thr Asn Gly Arg Cys Ile Thr Leu		
35	40	45
Leu Trp Lys Cys Asp Gly Asp Glu Asp Cys Thr Asp Gly Ser Asp Glu		
50	55	60
Lys Asn Cys Val Lys Lys Thr Cys Ala Glu Ser Asp Phe Val Cys Lys		
65	70	75
Asn Gly Gln Cys Val Pro Asn Arg Trp Gln Cys Asp Gly Asp Pro Asp		
85	90	95
Cys Glu Asp Gly Ser Asp Glu Ser Pro Glu Gln Cys His Met Arg Thr		
100	105	110
Cys Arg Ile Asn Glu Ile Ser Cys Gly Ala Arg Ser Thr Gln Cys Ile		
115	120	125
Pro Glu Ser Trp Arg Cys Asp Gly Glu Asn Asp Cys Asp Asn Gly Glu		
130	135	140
Asp Glu Glu Asn Cys Gly Asn Ile Thr Cys Ser Ala Asp Glu Phe Thr		
145	150	155
Cys Ser Ser Gly Arg Cys Val Ser Arg Asn Phe Val Cys Asn Gly Gln		
165	170	175
Asp Asp Cys Asp Asp Gly Ser Asp Glu Leu Asp Cys Ala Pro Pro Thr		
180	185	190
Cys Gly Ala His Glu Phe Gln Cys Arg Thr Ser Ser Cys Ile Pro Leu		
195	200	205

Ser Trp Val Cys Asp Asp Asp Ala Asp Cys Ser Asp Gln Ser Asp Glu  
 210 215 220  
 Ser Leu Glu Gln Cys Gly Arg Gln Pro Val Ile His Thr Lys Cys Pro  
 225 230 235 240  
 Thr Ser Glu Ile Gln Cys Gly Ser Gly Glu Cys Ile His Lys Lys Trp  
 245 250 255  
 Arg Cys Asp Gly Asp Pro Asp Cys Lys Asp Gly Ser Asp Glu Val Asn  
 260 265 270  
 Cys Pro Ser Arg Thr Cys Arg Pro Asp Gln Phe Glu Cys Glu Asp Gly  
 275 280 285  
 Ser Cys Ile His Gly Ser Arg Gln Cys Asn Gly Ile Arg Asp Cys Val  
 290 295 300  
 Asp Gly Ser Asp Glu Val Asn Cys Lys Asn Val Asn Gln Cys Leu Gly  
 305 310 315 320  
 Pro Gly Lys Phe Lys Cys Arg Ser Gly Glu Cys Ile Asp Ile Thr Lys  
 325 330 335  
 Val Cys Asp Gln Glu Gln Asp Cys Arg Asp Trp Ser Asp Glu Pro Leu  
 340 345 350  
 Lys Glu Cys His Ile Asn Glu Cys Leu Val Asn Asn Gly Gly Cys Ser  
 355 360 365  
 His Ile Cys Lys Asp Leu Val Ile Gly Tyr Glu Cys Asp Cys Ala Ala  
 370 375 380  
 Gly Phe Glu Leu Ile Asp Arg Lys Thr Cys Gly Asp Ile Asp Glu Cys  
 385 390 395 400  
 Gln Asn Pro Gly Ile Cys Ser Gln Ile Cys Ile Asn Leu Lys Gly Gly  
 405 410 415  
 Tyr Lys Cys Glu Cys Ser Arg Gly Tyr Gln Met Asp Leu Ala Thr Gly  
 420 425 430  
 Val Cys Lys Ala Val Gly Lys Glu Pro Ser Leu Ile Phe Thr Asn Arg  
 435 440 445  
 Arg Asp Ile Arg Lys Ile Gly Leu Glu Arg Lys Glu Tyr Ile Gln Leu  
 450 455 460  
 Val Glu Gln Leu Arg Asn Thr Val Ala Leu Asp Ala Asp Ile Ala Ala  
 465 470 475 480  
 Gln Lys Leu Phe Trp Ala Asp Leu Ser Gln Lys Ala Ile Phe Ser Ala  
 485 490 495  
 Ser Ile Asp Asp Lys Val Gly Arg His Phe Lys Met Ile Asp Asn Val  
 500 505 510  
 Tyr Asn Pro Ala Ala Ile Ala Val Asp Trp Val Tyr Lys Thr Ile Tyr  
 515 520 525  
 Trp Thr Asp Ala Ala Ser Lys Thr Ile Ser Val Ala Thr Leu Asp Gly  
 530 535 540  
 Thr Lys Arg Lys Phe Leu Phe Asn Ser Asp Leu Arg Glu Pro Ala Ser  
 545 550 555 560  
 Ile Ala Val Asp Pro Leu Ser Gly Phe Val Tyr Trp Ser Asp Trp Gly  
 565 570 575  
 Glu Pro Ala Lys Ile Glu Lys Ala Gly Met Asn Gly Phe Asp Arg Arg  
 580 585 590  
 Pro Leu Val Thr Glu Asp Ile Gln Trp Pro Asn Gly Ile Thr Leu Asp  
 595 600 605  
 Leu Val Lys Ser Arg Leu Tyr Trp Leu Asp Ser Lys Leu His Met Leu  
 610 615 620  
 Ser Ser Val Asp Leu Asn Gly Gln Asp Arg Arg Ile Val Leu Lys Ser  
 625 630 635 640  
 Leu Glu Phe Leu Ala His Pro Leu Ala Leu Thr Ile Phe Glu Asp Arg  
 645 650 655  
 Val Tyr Trp Ile Asp Gly Glu Asn Glu Ala Val Tyr Gly Ala Asn Lys  
 660 665 670  
 Phe Thr Gly Ser Glu Leu Ala Thr Leu Val Asn Asn Leu Asn Asp Ala

675	680	685
Gln Asp Ile Ile Ile Tyr His Glu Leu Val Gln Pro Ser Gly Lys Asn		
690	695	700
Trp Cys Glu Glu Asp Met Glu Asn Gly Gly Cys Glu Tyr Leu Cys Leu		
705	710	715
Pro Ala Pro Gln Ile Asn Asp His Ser Pro Lys Tyr Thr Cys Ser Cys		
725	730	735
Pro Asn Gly Tyr Asn Leu Glu Glu Asn Gly Arg Glu Cys Gln Ser Thr		
740	745	750
Ser Thr Pro Val Thr Tyr Ser Glu Thr Lys Asp Val Asn Thr Thr Asp		
755	760	765
Ile Leu Arg Thr Ser Gly Leu Val Pro Gly Gly Ile Asn Val Thr Thr		
770	775	780
Ala Val Ser Glu Val Ser Val Pro Pro Lys Gly Thr Ser Ala Ala Trp		
785	790	795
Ala Ile Leu Pro Leu Leu Leu Val Met Ala Ala Val Gly Gly Tyr		
805	810	815
Leu Met Trp Arg Asn Trp Gln His Lys Asn Met Lys Ser Met Asn Phe		
820	825	830
Asp Asn Pro Val Tyr Leu Lys Thr Thr Glu Glu Asp Leu Ser Ile Asp		
835	840	845
Ile Gly Arg His Ser Ala Ser Val Gly His Thr Tyr Pro Ala Ile Ser		
850	855	860
Val Val Ser Thr Asp Asp Asp Leu Ala		
865	870	

<210> 11  
 <211> 1280  
 <212> PRT  
 <213> Homo sapiens

<400> 11		
Met Asp Leu Glu Gly Asp Arg Asn Gly Gly Ala Lys Lys Lys Asn Phe		
1	5	10
Phe Lys Leu Asn Asn Lys Ser Glu Lys Asp Lys Lys Glu Lys Lys Pro		
20	25	30
Thr Val Ser Val Phe Ser Met Phe Arg Tyr Ser Asn Trp Leu Asp Lys		
35	40	45
Leu Tyr Met Val Val Gly Thr Leu Ala Ala Ile Ile His Gly Ala Gly		
50	55	60
Leu Pro Leu Met Met Leu Val Phe Gly Glu Met Thr Asp Ile Phe Ala		
65	70	75
Asn Ala Gly Asn Leu Glu Asp Leu Met Ser Asn Ile Thr Asn Arg Ser		
85	90	95
Asp Ile Asn Asp Thr Gly Phe Phe Met Asn Leu Glu Glu Asp Met Thr		
100	105	110
Arg Tyr Ala Tyr Tyr Tyr Ser Gly Ile Gly Ala Gly Val Leu Val Ala		
115	120	125
Ala Tyr Ile Gln Val Ser Phe Trp Cys Leu Ala Ala Gly Arg Gln Ile		
130	135	140
His Lys Ile Arg Lys Gln Phe Phe His Ala Ile Met Arg Gln Glu Ile		
145	150	155
Gly Trp Phe Asp Val His Asp Val Gly Glu Leu Asn Thr Arg Leu Thr		
165	170	175
Asp Asp Val Ser Lys Ile Asn Glu Gly Ile Gly Asp Lys Ile Gly Met		
180	185	190
Phe Phe Gln Ser Met Ala Thr Phe Phe Thr Gly Phe Ile Val Gly Phe		
195	200	205

Thr Arg Gly Trp Lys Leu Thr Leu Val Ile Leu Ala Ile Ser Pro Val  
 210 215 220  
 Leu Gly Leu Ser Ala Ala Val Trp Ala Lys Ile Leu Ser Ser Phe Thr  
 225 230 235 240  
 Asp Lys Glu Leu Leu Ala Tyr Ala Lys Ala Gly Ala Val Ala Glu Glu  
 245 250 255  
 Val Leu Ala Ala Ile Arg Thr Val Ile Ala Phe Gly Gly Gln Lys Lys  
 260 265 270  
 Glu Leu Glu Arg Tyr Asn Lys Asn Leu Glu Glu Ala Lys Arg Ile Gly  
 275 280 285  
 Ile Lys Lys Ala Ile Thr Ala Asn Ile Ser Ile Gly Ala Ala Phe Leu  
 290 295 300  
 Leu Ile Tyr Ala Ser Tyr Ala Leu Ala Phe Trp Tyr Gly Thr Thr Leu  
 305 310 315 320  
 Val Leu Ser Gly Glu Tyr Ser Ile Gly Gln Val Leu Thr Val Phe Phe  
 325 330 335  
 Ser Val Leu Ile Gly Ala Phe Ser Val Gly Gln Ala Ser Pro Ser Ile  
 340 345 350  
 Glu Ala Phe Ala Asn Ala Arg Gly Ala Ala Tyr Glu Ile Phe Lys Ile  
 355 360 365  
 Ile Asp Asn Lys Pro Ser Ile Asp Ser Tyr Ser Lys Ser Gly His Lys  
 370 375 380  
 Pro Asp Asn Ile Lys Gly Asn Leu Glu Phe Arg Asn Val His Phe Ser  
 385 390 395 400  
 Tyr Pro Ser Arg Lys Glu Val Lys Ile Leu Lys Gly Leu Asn Leu Lys  
 405 410 415  
 Val Gln Ser Gly Gln Thr Val Ala Leu Val Gly Asn Ser Gly Cys Gly  
 420 425 430  
 Lys Ser Thr Thr Val Gln Leu Met Gln Arg Leu Tyr Asp Pro Thr Glu  
 435 440 445  
 Gly Met Val Ser Val Asp Gly Gln Asp Ile Arg Thr Ile Asn Val Arg  
 450 455 460  
 Phe Leu Arg Glu Ile Ile Gly Val Val Ser Gln Glu Pro Val Leu Phe  
 465 470 475 480  
 Ala Thr Thr Ile Ala Glu Asn Ile Arg Tyr Gly Arg Glu Asn Val Thr  
 485 490 495  
 Met Asp Glu Ile Glu Lys Ala Val Lys Glu Ala Asn Ala Tyr Asp Phe  
 500 505 510  
 Ile Met Lys Leu Pro His Lys Phe Asp Thr Leu Val Gly Glu Arg Gly  
 515 520 525  
 Ala Gln Leu Ser Gly Gly Gln Lys Gln Arg Ile Ala Ile Ala Arg Ala  
 530 535 540  
 Leu Val Arg Asn Pro Lys Ile Leu Leu Asp Glu Ala Thr Ser Ala  
 545 550 555 560  
 Leu Asp Thr Glu Ser Glu Ala Val Val Gln Val Ala Leu Asp Lys Ala  
 565 570 575  
 Arg Lys Gly Arg Thr Thr Ile Val Ile Ala His Arg Leu Ser Thr Val  
 580 585 590  
 Arg Asn Ala Asp Val Ile Ala Gly Phe Asp Asp Gly Val Ile Val Glu  
 595 600 605  
 Lys Gly Asn His Asp Glu Leu Met Lys Glu Lys Gly Ile Tyr Phe Lys  
 610 615 620  
 Leu Val Thr Met Gln Thr Ala Gly Asn Glu Val Glu Leu Glu Asn Ala  
 625 630 635 640  
 Ala Asp Glu Ser Lys Ser Glu Ile Asp Ala Leu Glu Met Ser Ser Asn  
 645 650 655  
 Asp Ser Arg Ser Ser Leu Ile Arg Lys Arg Ser Thr Arg Arg Ser Val  
 660 665 670  
 Arg Gly Ser Gln Ala Gln Asp Arg Lys Leu Ser Thr Lys Glu Ala Leu

675	680	685
Asp Glu Ser Ile Pro Pro Val Ser Phe Trp Arg	Ile Met Lys Leu Asn	
690	695	700
Leu Thr Glu Trp Pro Tyr Phe Val Val Gly Val	Phe Cys Ala Ile Ile	
705	710	715
Asn Gly Gly Leu Gln Pro Ala Phe Ala Ile Ile	Phe Ser Lys Ile Ile	
725	730	735
Gly Val Phe Thr Arg Ile Asp Asp Pro Glu Thr Lys Arg	Gln Asn Ser	
740	745	750
Asn Leu Phe Ser Leu Leu Phe Leu Ala Leu Gly	Ile Ile Ser Phe Ile	
755	760	765
Thr Phe Phe Leu Gln Gly Phe Thr Phe Gly Lys	Ala Gly Glu Ile Leu	
770	775	780
Thr Lys Arg Leu Arg Tyr Met Val Phe Arg Ser	Met Leu Arg Gln Asp	
785	790	795
Val Ser Trp Phe Asp Asp Pro Lys Asn Thr Thr	Gly Ala Leu Thr Thr	
805	810	815
Arg Leu Ala Asn Asp Ala Ala Gln Val Lys Gly	Ala Ile Gly Ser Arg	
820	825	830
Leu Ala Val Ile Thr Gln Asn Ile Ala Asn Leu Gly	Thr Gly Ile Ile	
835	840	845
Ile Ser Phe Ile Tyr Gly Trp Gln Leu Thr Leu	Leu Leu Ala Ile	
850	855	860
Val Pro Ile Ile Ala Ile Ala Gly Val Val Glu	Met Lys Met Leu Ser	
865	870	875
Gly Gln Ala Leu Lys Asp Lys Glu Leu Glu Gly	Ala Gly Lys Ile	
885	890	895
Ala Thr Glu Ala Ile Glu Asn Phe Arg Thr Val Val	Ser Leu Thr Gln	
900	905	910
Glu Gln Lys Phe Glu His Met Tyr Ala Gln Ser Leu	Gln Val Pro Tyr	
915	920	925
Arg Asn Ser Leu Arg Lys Ala His Ile Phe Gly	Ile Thr Phe Ser Phe	
930	935	940
Thr Gln Ala Met Met Tyr Phe Ser Tyr Ala Gly	Cys Phe Arg Phe Gly	
945	950	955
Ala Tyr Leu Val Ala His Lys Leu Met Ser Phe	Glu Asp Val Leu Leu	
965	970	975
Val Phe Ser Ala Val Val Phe Gly Ala Met Ala Val	Gly Gln Val Ser	
980	985	990
Ser Phe Ala Pro Asp Tyr Ala Lys Ala Lys Ile	Ser Ala Ala His Ile	
995	1000	1005
Ile Met Ile Ile Glu Lys Thr Pro Leu Ile Asp	Ser Tyr Ser Thr Glu	
1010	1015	1020
Gly Leu Met Pro Asn Thr Leu Glu Gly Asn Val	Thr Phe Gly Glu Val	
1025	1030	1035
Val Phe Asn Tyr Pro Thr Arg Pro Asp Ile Pro	Val Leu Gln Gly Leu	
1045	1050	1055
Ser Leu Glu Val Lys Lys Gly Gln Thr Leu Ala	Leu Val Gly Ser Ser	
1060	1065	1070
Gly Cys Gly Lys Ser Thr Val Val Gln Leu Leu	Glu Arg Phe Tyr Asp	
1075	1080	1085
Pro Leu Ala Gly Lys Val Leu Leu Asp Gly Lys	Glu Ile Lys Arg Leu	
1090	1095	1100
Asn Val Gln Trp Leu Arg Ala His Leu Gly	Ile Val Ser Gln Glu Pro	
1105	1110	1115
Ile Leu Phe Asp Cys Ser Ile Ala Glu Asn Ile	Ala Tyr Gly Asp Asn	
1125	1130	1135
Ser Arg Val Val Ser Gln Glu Glu Ile Val Arg	Ala Ala Lys Glu Ala	
1140	1145	1150

Asn Ile His Ala Phe Ile Glu Ser Leu Pro Asn Lys Tyr Ser Thr Lys			
1155	1160	1165	
Val Gly Asp Lys Gly Thr Gln Leu Ser Gly Gly Gln Lys Gln Arg Ile			
1170	1175	1180	
Ala Ile Ala Arg Ala Leu Val Arg Gln Pro His Ile Leu Leu Leu Asp			
1185	1190	1195	1200
Glu Ala Thr Ser Ala Leu Asp Thr Glu Ser Glu Lys Val Val Gln Glu			
1205	1210	1215	
Ala Leu Asp Lys Ala Arg Glu Gly Arg Thr Cys Ile Val Ile Ala His			
1220	1225	1230	
Arg Leu Ser Thr Ile Gln Asn Ala Asp Leu Ile Val Val Phe Gln Asn			
1235	1240	1245	
Gly Arg Val Lys Glu His Gly Thr His Gln Gln Leu Leu Ala Gln Lys			
1250	1255	1260	
Gly Ile Tyr Phe Ser Met Val Ser Val Gln Ala Gly Thr Lys Arg Gln			
1265	1270	1275	1280

<210> 12  
 <211> 1276  
 <212> PRT  
 <213> Mus musculus

<400> 12			
Met Glu Phe Glu Glu Asn Leu Lys Gly Arg Ala Asp Lys Asn Phe Ser			
1	5	10	15
Lys Met Gly Lys Lys Ser Lys Lys Glu Lys Lys Glu Lys Lys Pro Ala			
20	25	30	
Val Gly Val Phe Gly Met Phe Arg Tyr Ala Asp Trp Leu Asp Lys Leu			
35	40	45	
Cys Met Ile Leu Gly Thr Leu Ala Ala Ile Ile His Gly Thr Leu Leu			
50	55	60	
Pro Leu Leu Met Leu Val Phe Gly Asn Met Thr Asp Ser Phe Thr Lys			
65	70	75	80
Ala Glu Ala Ser Ile Leu Pro Ser Ile Thr Asn Gln Ser Gly Pro Asn			
85	90	95	
Ser Thr Leu Ile Ile Ser Asn Ser Ser Leu Glu Glu Glu Met Ala Ile			
100	105	110	
Tyr Ala Tyr Tyr Tyr Gly Ile Gly Ala Gly Val Leu Ile Val Ala			
115	120	125	
Tyr Ile Gln Val Ser Leu Trp Cys Leu Ala Ala Gly Arg Gln Ile His			
130	135	140	
Lys Ile Arg Gln Lys Phe Phe His Ala Ile Met Asn Gln Glu Ile Gly			
145	150	155	160
Trp Phe Asp Val His Asp Val Gly Glu Leu Asn Thr Arg Lys Thr Asp			
165	170	175	
Asp Val Ser Lys Ile Asn Asp Gly Ile Gly Asp Lys Ile Gly Met Phe			
180	185	190	
Phe Gln Ser Ile Thr Thr Phe Leu Ala Gly Phe Ile Ile Gly Phe Ile			
195	200	205	
Ser Gly Trp Lys Leu Thr Leu Val Ile Leu Ala Val Ser Pro Leu Ile			
210	215	220	
Gly Leu Ser Ser Ala Leu Trp Ala Lys Val Leu Thr Ser Phe Thr Asn			
225	230	235	240
Lys Glu Leu Gln Ala Tyr Ala Lys Ala Gly Ala Val Ala Glu Glu Val			
245	250	255	
Leu Ala Ala Ile Arg Thr Val Ile Ala Phe Gly Gly Gln Gln Lys Glu			
260	265	270	
Leu Glu Arg Tyr Asn Lys Asn Leu Glu Glu Ala Lys Asn Val Gly Ile			

275	280	285													
Lys	Lys	Ala	Ile	Thr	Ala	Ser	Ile	Ser	Ile	Gly	Ile	Ala	Tyr	Leu	Leu
290							295				300				
Val	Tyr	Ala	Ser	Tyr	Ala	Leu	Ala	Phe	Trp	Tyr	Gly	Thr	Ser	Leu	Val
305							310				315				320
Leu	Ser	Asn	Glu	Tyr	Ser	Ile	Gly	Glu	Val	Leu	Thr	Val	Phe	Phe	Ser
							325				330				335
Ile	Leu	Leu	Gly	Thr	Phe	Ser	Ile	Gly	His	Leu	Ala	Pro	Asn	Ile	Glu
							340				345				350
Ala	Phe	Ala	Asn	Ala	Arg	Gly	Ala	Ala	Phe	Glu	Ile	Phe	Lys	Ile	Ile
							355				360				365
Asp	Asn	Glu	Pro	Ser	Ile	Asp	Ser	Phe	Ser	Thr	Lys	Gly	Tyr	Lys	Pro
							370				375				380
Asp	Ser	Ile	Met	Gly	Asn	Leu	Glu	Phe	Lys	Asn	Val	His	Phe	Asn	Tyr
							385				390				400
Pro	Ser	Arg	Ser	Glu	Val	Gln	Ile	Leu	Lys	Gly	Leu	Asn	Leu	Lys	Val
							405				410				415
Lys	Ser	Gly	Gln	Thr	Val	Ala	Leu	Val	Gly	Asn	Ser	Gly	Cys	Gly	Lys
							420				425				430
Ser	Thr	Thr	Val	Gln	Leu	Met	Gln	Arg	Leu	Tyr	Asp	Pro	Leu	Glu	Gly
							435				440				445
Val	Val	Ser	Ile	Asp	Gly	Gln	Asp	Ile	Arg	Thr	Ile	Asn	Val	Arg	Tyr
							450				455				460
Leu	Arg	Glu	Ile	Ile	Gly	Val	Val	Ser	Gln	Glu	Pro	Val	Leu	Phe	Ala
							465				470				480
Thr	Thr	Ile	Ala	Glu	Asn	Ile	Arg	Tyr	Gly	Arg	Glu	Asp	Val	Thr	Met
							485				490				495
Asp	Glu	Ile	Glu	Lys	Ala	Val	Lys	Glu	Ala	Asn	Ala	Tyr	Asp	Phe	Ile
							500				505				510
Met	Lys	Leu	Pro	His	Gln	Phe	Asp	Thr	Leu	Val	Gly	Glu	Arg	Gly	Ala
							515				520				525
Gln	Leu	Ser	Gly	Gly	Gln	Lys	Gln	Arg	Ile	Ala	Ile	Ala	Arg	Ala	Leu
							530				535				540
Val	Arg	Asn	Pro	Lys	Ile	Leu	Leu	Leu	Asp	Glu	Ala	Thr	Ser	Ala	Leu
							545				550				560
Asp	Thr	Glu	Ser	Glu	Ala	Val	Val	Gln	Ala	Ala	Leu	Asp	Lys	Ala	Arg
							565				570				575
Glu	Gly	Arg	Thr	Thr	Ile	Val	Ile	Ala	His	Arg	Leu	Ser	Thr	Val	Arg
							580				585				590
Asn	Ala	Asp	Val	Ile	Ala	Gly	Phe	Asp	Gly	Gly	Val	Ile	Val	Glu	Gln
							595				600				605
Gly	Asn	His	Asp	Glu	Leu	Met	Arg	Glu	Lys	Gly	Ile	Tyr	Phe	Lys	Leu
							610				615				620
Val	Met	Thr	Gln	Thr	Arg	Gly	Asn	Glu	Ile	Glu	Pro	Gly	Asn	Asn	Ala
							625				630				640
Tyr	Gly	Ser	Gln	Ser	Asp	Thr	Asp	Ala	Ser	Glu	Leu	Thr	Ser	Glu	Glu
							645				650				655
Ser	Lys	Ser	Pro	Leu	Ile	Arg	Arg	Ser	Ile	Tyr	Arg	Ser	Val	His	Arg
							660				665				670
Lys	Gln	Asp	Gln	Glu	Arg	Arg	Leu	Ser	Met	Lys	Glu	Ala	Val	Asp	Glu
							675				680				685
Asp	Val	Pro	Leu	Val	Ser	Phe	Trp	Arg	Ile	Leu	Asn	Leu	Asn	Leu	Ser
							690				695				700
Glu	Trp	Pro	Tyr	Leu	Leu	Val	Gly	Val	Leu	Cys	Ala	Val	Ile	Asn	Gly
							705				710				720
Cys	Ile	Gln	Pro	Val	Phe	Ala	Ile	Val	Phe	Ser	Arg	Ile	Val	Gly	Val
							725				730				735
Phe	Ser	Arg	Asp	Asp	Asp	His	Glu	Thr	Lys	Arg	Gln	Asn	Cys	Asn	Leu
							740				745				750

Phe Ser Leu Phe Phe Leu Val Met Gly Leu Ile Ser Phe Val Thr Tyr  
 755 760 765  
 Phe Phe Gln Gly Phe Thr Phe Gly Lys Ala Gly Glu Ile Leu Thr Lys  
 770 775 780  
 Arg Val Arg Tyr Met Val Phe Lys Ser Met Leu Arg Gln Asp Ile Ser  
 785 790 795 800  
 Trp Phe Asp Asp His Lys Asn Ser Thr Gly Ser Leu Thr Thr Arg Leu  
 805 810 815  
 Ala Ser Asp Ala Ser Ser Val Lys Gly Ala Met Gly Ala Arg Leu Ala  
 820 825 830  
 Val Val Thr Gln Asn Val Ala Asn Leu Gly Thr Gly Val Ile Leu Ser  
 835 840 845  
 Leu Val Tyr Gly Trp Gln Leu Thr Leu Leu Val Val Ile Ile Pro  
 850 855 860  
 Leu Ile Val Leu Gly Gly Ile Ile Glu Met Lys Leu Leu Ser Gly Gln  
 865 870 875 880  
 Ala Leu Lys Asp Lys Lys Gln Leu Glu Ile Ser Gly Lys Ile Ala Thr  
 885 890 895  
 Glu Ala Ile Glu Asn Phe Arg Thr Ile Val Ser Leu Thr Arg Glu Gln  
 900 905 910  
 Lys Phe Glu Thr Met Tyr Ala Gln Ser Leu Gln Val Pro Tyr Arg Asn  
 915 920 925  
 Ala Met Lys Ala His Val Phe Gly Ile Thr Phe Ser Phe Thr Gln  
 930 935 940  
 Ala Met Met Tyr Phe Ser Tyr Ala Ala Cys Phe Arg Phe Gly Ala Tyr  
 945 950 955 960  
 Leu Val Ala Gln Gln Leu Met Thr Phe Glu Asn Val Met Leu Val Phe  
 965 970 975  
 Ser Ala Val Val Phe Gly Ala Met Ala Ala Gly Asn Thr Ser Ser Phe  
 980 985 990  
 Ala Pro Asp Tyr Ala Lys Ala Lys Val Ser Ala Ser His Ile Ile Arg  
 995 1000 1005  
 Ile Ile Glu Lys Thr Pro Glu Ile Asp Ser Tyr Ser Thr Glu Gly Leu  
 1010 1015 1020  
 Lys Pro Thr Leu Leu Glu Gly Asn Val Lys Phe Asn Gly Val Gln Phe  
 1025 1030 1035 1040  
 Asn Tyr Pro Thr Arg Pro Asn Ile Pro Val Leu Gln Gly Leu Ser Leu  
 1045 1050 1055  
 Glu Val Lys Lys Gly Gln Thr Leu Ala Leu Val Gly Ser Ser Gly Cys  
 1060 1065 1070  
 Gly Lys Ser Thr Val Val Gln Leu Leu Glu Arg Phe Tyr Asp Pro Met  
 1075 1080 1085  
 Ala Gly Ser Val Phe Leu Asp Gly Lys Glu Ile Lys Gln Leu Asn Val  
 1090 1095 1100  
 Gln Trp Leu Arg Ala His Leu Gly Ile Val Ser Gln Glu Pro Ile Leu  
 1105 1110 1115 1120  
 Phe Asp Cys Ser Ile Ala Glu Asn Ile Ala Tyr Gly Asp Asn Ser Arg  
 1125 1130 1135  
 Ala Val Ser His Glu Glu Ile Val Arg Ala Ala Lys Glu Ala Asn Ile  
 1140 1145 1150  
 His Gln Phe Ile Asp Ser Leu Pro Asp Lys Tyr Asn Thr Arg Val Gly  
 1155 1160 1165  
 Asp Lys Gly Thr Gln Leu Ser Gly Gly Gln Lys Gln Arg Ile Ala Ile  
 1170 1175 1180  
 Ala Arg Ala Leu Val Arg Gln Pro His Ile Leu Leu Asp Glu Ala  
 1185 1190 1195 1200  
 Thr Ser Ala Leu Asp Thr Glu Ser Glu Lys Val Val Gln Glu Ala Leu  
 1205 1210 1215  
 Asp Lys Ala Arg Glu Gly Arg Thr Cys Ile Val Ile Ala His Arg Leu

1220 1225 1230  
Ser Thr Ile Gln Asn Ala Asp Leu Ile Val Val Ile Glu Asn Gly Lys  
1235 1240 1245  
Val Lys Glu His Gly Thr His Gln Gln Leu Leu Ala Gln Lys Gly Ile  
1250 1255 1260  
Tyr Phe Ser Met Val Gln Ala Gly Ala Lys Arg Ser  
1265 1270 1275